RESULT 1
LOL4_HUMAN
ID   LOL4_HUMAN   STANDARD;   PRT;   756 AA.
AC   Q96JB6; Q96DY1; Q96PC0; Q9H6T5;
DT   28-FEB-2003 (Rel. 41, Created)
DT   28-FEB-2003 (Rel. 41, Last sequence update)
DT   10-OCT-2003 (Rel. 42, Last annotation update)
DE   Lysyl oxidase homolog 4 precursor (EC 1.4.3.-) (Lysyl oxidase-like
DE   protein 4) (Lysyl oxidase related protein C).
GN   LOXL4 OR LOXC.
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Testis;
RX   MEDLINE=21316447; PubMed=11292829;
RA   Ito H., Akiyama H., Iguchi H., Iyama K., Miyamoto M., Ohsawa K.,
RA   Nakamura T.;
RT   "Molecular cloning and biological activity of a novel lysyl oxidase-
RT   related gene expressed in cartilage.";
RL   J. Biol. Chem. 276:24023-24029(2001).
RN   [2]
RP   SEQUENCE FROM N.A.
RX   MEDLINE=21550107; PubMed=11691589;
RA   Maeki J.M., Tikkanen H., Kivirikko K.I.;
RT   "Cloning and characterization of a fifth human lysyl oxidase
RT   isoenzyme: the third member of the lysyl oxidase-related subfamily
RT   with four scavenger receptor cysteine-rich domains.";
RL   Matrix Biol. 20:493-496(2001).
RN   [3]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Placenta;
RX   MEDLINE=21550106; PubMed=11691588;
RA   Asuncion L., Fogelgren B., Fong K.S.K., Fong S.F.T., Kim Y.,
RA   Csiszar K.;
RT   "A novel human lysyl oxidase-like gene (LOXL4) on chromosome 10q24 has
RT   an altered scavenger receptor cysteine rich domain.";
RL   Matrix Biol. 20:487-491(2001).
RN   [4]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Eye;
RX   MEDLINE=22388257; PubMed=12477932;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

```
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA   Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
RA   Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length
RT   human and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [5]
RP   SEQUENCE OF 492-756 FROM N.A.
RA   Kawabata A., Hikiji T., Kobatake N., Inagaki H., Ikema Y., Okamoto S.,
RA   Okitani R., Ota T., Suzuki Y., Obayashi M., Nishi T., Shibahara T.,
RA   Tanaka T., Nakamura Y., Isogai T., Sugano S.;
RT   "NEDO human cDNA sequencing project.";
RL   Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
CC   -!- FUNCTION: May modulate the formation of a collagenous
CC       extracellular matrix.
CC   -!- COFACTOR: Copper and LTQ (By similarity).
CC   -!- SUBCELLULAR LOCATION: Extracellular (Potential).
CC   -!- TISSUE SPECIFICITY: Expressed in many tissues, the highest levels
CC       among the tissues studied being in the skeletal muscle, testis and
CC       pancreas. Expressed in cartilage.
CC   -!- PTM: The lysine tyrosylquinone cross-link (LTQ) is generated by
CC       condensation of the epsilon-amino group of a lysine with a
CC       topaquinone produced by oxidation of tyrosine.
CC   -!- SIMILARITY: Contains 4 SRCR domains.
CC   -!- SIMILARITY: Belongs to the lysyl oxidase family.
CC   ----------------------------------------------------------------------
CC   This SWISS-PROT entry is copyright. It is produced through a collaboration
CC   between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC   the European Bioinformatics Institute. There are no restrictions on its
CC   use by non-profit institutions as long as its content is in no way
CC   modified and this statement is not removed. Usage by and for commercial
CC   entities requires a license agreement (See http://www.isb-sib.ch/announce/
CC   or send an email to license@isb-sib.ch).
CC   ----------------------------------------------------------------------
DR   EMBL; AF338441; AAK71934.1; -.
DR   EMBL; AY036093; AAK64186.1; -.
DR   EMBL; AF395336; AAL27543.1; -.
DR   EMBL; BC007522; AAH07522.1; ALT_INIT.
DR   EMBL; BC013153; AAH13153.1; -.
DR   EMBL; AK025542; BAB15167.1; -.
DR   Genew; HGNC:17171; LOXL4.
DR   MIM; 607318; -.
DR   InterPro; IPR001695; Lysyl_oxidase.
DR   InterPro; IPR001190; Srcr_receptor.
DR   Pfam; PF01186; Lysyl_oxidase; 1.
DR   Pfam; PF00530; SRCR; 4.
DR   PRINTS; PR00074; LYSYLOXIDASE.
DR   PRINTS; PR00258; SPERACTRCPTR.
DR   ProDom; PD013887; Lysyl_oxidase; 1.
DR   SMART; SM00202; SR; 4.
DR   PROSITE; PS00926; LYSYL_OXIDASE; FALSE_NEG.
DR   PROSITE; PS00420; SRCR_1; 1.
DR   PROSITE; PS50287; SRCR_2; 4.
KW   Oxidoreductase; Copper; Glycoprotein; Repeat; Signal; LTQ.
FT   SIGNAL        1     24       POTENTIAL.
FT   CHAIN        25    756       LYSYL OXIDASE HOMOLOG 4.
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FT   DOMAIN       32    133       SRCR 1.
FT   DOMAIN      159    287       SRCR 2.
FT   DOMAIN      311    411       SRCR 3.
FT   DOMAIN      421    529       SRCR 4.
FT   DOMAIN      533    736       LYSYL-OXIDASE LIKE.
FT   METAL       611    611       COPPER (POTENTIAL).
FT   METAL       613    613       COPPER (POTENTIAL).
FT   METAL       615    615       COPPER (POTENTIAL).
FT   CROSSLNK    638    674       Lysine tyrosylquinone (Lys-Tyr)
FT                                (By similarity).
FT   MOD_RES     674    674       TOPAQUINONE (BY SIMILARITY).
FT   CARBOHYD    198    198       N-LINKED (GLCNAC. . .) (POTENTIAL).
FT   CARBOHYD    629    629       N-LINKED (GLCNAC. . .) (POTENTIAL).
FT   CONFLICT      3      3       W -> R (IN REF. 4).
FT   CONFLICT    101    101       R -> Q (IN REF. 4).
FT   CONFLICT    405    405       D -> A (IN REF. 4).
FT   CONFLICT    493    493       S -> G (IN REF. 3).
FT   CONFLICT    539    539       A -> T (IN REF. 3).
FT   CONFLICT    542    542       V -> A (IN REF. 3).
FT   CONFLICT    703    703       Y -> H (IN REF. 3).
SQ   SEQUENCE   756 AA;  84483 MW;  13051ACADB922BBC CRC64;

  Query Match             99.9%;  Score 4174;  DB 1;  Length 756;
  Best Local Similarity   99.9%;  Pred. No. 0;
  Matches  755;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MAWSPPATLFLFLLLLGQPPPSRPQSLGTTKLRLVGPESKPEEGRLEVLHQGQWGTVCDD 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MAWSPPATLFLFLLLLGQPPPSRPQSLGTTKLRLVGPESKPEEGRLEVLHQGQWGTVCDD 60

Qy         61 NFAIQEATVACRQLGFEAALTWAHSAKYGQGEGPIWLDNVRCVGTESSLDQCGSNGWGVS 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 NFAIQEATVACRQLGFEAALTWAHSAKYGQGEGPIWLDNVRCVGTESSLDQCGSNGWGVS 120

Qy        121 DCSHSEDVGVICHPRRHRGYLSETVSNALGPQGRRLEEVRLKPILASAKQHSPVTEGAVE 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 DCSHSEDVGVICHPRRHRGYLSETVSNALGPQGRRLEEVRLKPILASAKQHSPVTEGAVE 180

Qy        181 VKYEGHWRQVCDQGWTMNNSRVVCGMLGFPSEVPVDSHYYRKVWDLKMRDPKSRLKSLTN 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 VKYEGHWRQVCDQGWTMNNSRVVCGMLGFPSEVPVDSHYYRKVWDLKMRDPKSRLKSLTN 240

Qy        241 KNSFWIHQVTCLGTEPHMANCQVQVAPARGKLRPACPGGMHAVVSCVAGPHFRPPKTKPQ 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 KNSFWIHQVTCLGTEPHMANCQVQVAPARGKLRPACPGGMHAVVSCVAGPHFRPPKTKPQ 300

Qy        301 RKGSWAEEPRVRLRSGAQVGEGRVEVLMNRQWGTVCDHRWNLISASVVCRQLGFGSAREA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 RKGSWAEEPRVRLRSGAQVGEGRVEVLMNRQWGTVCDHRWNLISASVVCRQLGFGSAREA 360

Qy        361 LFGARLGQGLGPIHLSEVRCRGYERTLSDCPALEGSQNGCQHENAAAVRCNVPNMGFQNQ 420
              |||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db        361 LFGARLGQGLGPIHLSEVRCRGYERTLSDCPALEGSQNGCQHENDAAVRCNVPNMGFQNQ 420

Qy        421 VRLAGGRIPEEGLLEVQVEVNGVPRWGSVCSENWGLTEAMVACRQLGLGFAIHAYKETWF 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 VRLAGGRIPEEGLLEVQVEVNGVPRWGSVCSENWGLTEAMVACRQLGLGFAIHAYKETWF 480
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Qy        481 WSGTPRAQEVVMSGVRCSGTELALQQCQRHGPVHCSHGGGRFLAGVSCMDSAPDLVMNAQ 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        481 WSGTPRAQEVVMSGVRCSGTELALQQCQRHGPVHCSHGGGRFLAGVSCMDSAPDLVMNAQ 540

Qy        541 LVQETAYLEDRPLSQLYCAHEENCLSKSADHMDWPYGYRRLLRFSTQIYNLGRTDFRPKT 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        541 LVQETAYLEDRPLSQLYCAHEENCLSKSADHMDWPYGYRRLLRFSTQIYNLGRTDFRPKT 600

Qy        601 GRDSWVWHQCHRHYHSIEVFTHYDLLTLNGSKVAEGHKASFCLEDTNCPTGLQRRYACAN 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        601 GRDSWVWHQCHRHYHSIEVFTHYDLLTLNGSKVAEGHKASFCLEDTNCPTGLQRRYACAN 660

Qy        661 FGEQGVTVGCWDTYRHDIDCQWVDITDVGPGNYIFQVIVNPHYEVAESDFSNNMLQCRCK 720
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        661 FGEQGVTVGCWDTYRHDIDCQWVDITDVGPGNYIFQVIVNPHYEVAESDFSNNMLQCRCK 720

Qy        721 YDGHRVWLHNCHTGNSYPANAELSLEQEQRLRNNLI 756
              ||||||||||||||||||||||||||||||||||||
Db        721 YDGHRVWLHNCHTGNSYPANAELSLEQEQRLRNNLI 756
```

RESULT 2
AAM48743
ID AAM48743 standard; protein; 756 AA.
XX
AC AAM48743;
XX
DT 02-APR-2002 (first entry)
XX
DE Human 47765 lysyl oxidase SEQ ID NO 2.
XX
KW Human; 47765; lysyl oxidase; LSO; cytostatic; haemostatic; hepatotropic;
KW cardiant; osteopathic; dermatological; antiarteriosclerotic; vasotropic;
KW antiinflammatory; hypotensive; antiarrhythmic; cell proliferation;
KW growth; differentiation; leukaemia; tumour; cancer; bone; cartilage;
KW myeloproliferative; muscular; osteoporosis; cardiovascular; gene therapy;
KW chromosome mapping; tissue typing; forensic; pharmacogenomic; enzyme.
XX
OS Homo sapiens.
XX
PN WO200192495-A2.
XX
PD 06-DEC-2001.
XX
PF 29-MAY-2001; 2001WO-US017405.
XX
PR 26-MAY-2000; 2000US-0207650P.
XX
PA (MILL-) MILLENNIUM PHARM INC.
XX
PI Meyers R;
XX
DR WPI; 2002-122067/16.
DR N-PSDB; ABA96419, ABA96420.
XX
PT Novel human lysyl oxidase polypeptide, designated 47765, and
PT polynucleotides, useful in the diagnosis and treatment of cell
PT proliferation disorders, muscular disorders, bone disorders and skin
PT elasticity disorders.
XX
PS Claim 14; Fig 1; 115pp; English.
XX
CC The invention relates to human lysyl oxidase (LSO) polypeptide,
CC designated 47765 with cytostatic, haemostatic, hepatotropic, cardiant,
CC osteopathic, dermatological, antiarteriosclerotic, vasotropic,
CC antiinflammatory, hypotensive and antiarrhythmic activity. 47765
CC molecules are useful for identifying a compound which modulates the
CC activity of the protein, for developing novel diagnostic and therapeutic
CC agents for LSO-mediated or related disorders including cell
CC proliferation, growth or differentiation disorder (e.g. carcinoma,
CC leukaemia, tumour angiogenesis, hepatic disorders and haematopoietic,
CC myeloproliferative disorders), muscular disorders (e.g. cardiac muscle
CC disorder, paralysis, ataxia, myotonia and myokymia), bone disorders (e.g.
CC osteochondrosis and osteoporosis), skin elasticity disorders (e.g. cutis
CC laxa, Ehlers-Danlos type V syndrome), cardiovascular disorders (e.g.
CC arteriosclerosis, ischaemia reperfusion injury, restenosis, arterial
CC inflammation, vascular wall remodeling, tachycardia, vascular heart
CC disease, long QT syndrome, congestive heart failure, hypertension,

CC coronary artery disease and arrhythmia) or cartilage based disorders
CC (e.g. chondromalacia and polychondritis). The encoding polynucleotide is
CC useful in chromosome mapping, tissue typing, forensic identification, as
CC markers for pharmacogenomic profiling of a subject and in gene therapy
XX
SQ Sequence 756 AA;

Query Match 99.9%; Score 4174; DB 5; Length 756;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 755; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          1 MAWSPPATLFLFLLLLGQPPPSRPQSLGTTKLRLVGPESKPEEGRLEVLHQGQWGTVCDD 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MAWSPPATLFLFLLLLGQPPPSRPQSLGTTKLRLVGPESKPEEGRLEVLHQGQWGTVCDD 60

Qy         61 NFAIQEATVACRQLGFEAALTWAHSAKYGQGEGPIWLDNVRCVGTESSLDQCGSNGWGVS 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 NFAIQEATVACRQLGFEAALTWAHSAKYGQGEGPIWLDNVRCVGTESSLDQCGSNGWGVS 120

Qy        121 DCSHSEDVGVICHPRRHRGYLSETVSNALGPQGRRLEEVRLKPILASAKQHSPVTEGAVE 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 DCSHSEDVGVICHPRRHRGYLSETVSNALGPQGRRLEEVRLKPILASAKQHSPVTEGAVE 180

Qy        181 VKYEGHWRQVCDQGWTMNNSRVVCGMLGFPSEVPVDSHYYRKVWDLKMRDPKSRLKSLTN 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 VKYEGHWRQVCDQGWTMNNSRVVCGMLGFPSEVPVDSHYYRKVWDLKMRDPKSRLKSLTN 240

Qy        241 KNSFWIHQVTCLGTEPHMANCQVQVAPARGKLRPACPGGMHAVVSCVAGPHFRPPKTKPQ 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 KNSFWIHQVTCLGTEPHMANCQVQVAPARGKLRPACPGGMHAVVSCVAGPHFRPPKTKPQ 300

Qy        301 RKGSWAEEPRVRLRSGAQVGEGRVEVLMNRQWGTVCDHRWNLISASVVCRQLGFGSAREA 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 RKGSWAEEPRVRLRSGAQVGEGRVEVLMNRQWGTVCDHRWNLISASVVCRQLGFGSAREA 360

Qy        361 LFGARLGQGLGPIHLSEVRCRGYERTLSDCPALEGSQNGCQHENAAAVRCNVPNMGFQNQ 420
              |||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db        361 LFGARLGQGLGPIHLSEVRCRGYERTLSDCPALEGSQNGCQHENDAAVRCNVPNMGFQNQ 420

Qy        421 VRLAGGRIPEEGLLEVQVEVNGVPRWGSVCSENWGLTEAMVACRQLGLGFAIHAYKETWF 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 VRLAGGRIPEEGLLEVQVEVNGVPRWGSVCSENWGLTEAMVACRQLGLGFAIHAYKETWF 480

Qy        481 WSGTPRAQEVVMSGVRCSGTELALQQCQRHGPVHCSHGGGRFLAGVSCMDSAPDLVMNAQ 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        481 WSGTPRAQEVVMSGVRCSGTELALQQCQRHGPVHCSHGGGRFLAGVSCMDSAPDLVMNAQ 540

Qy        541 LVQETAYLEDRPLSQLYCAHEENCLSKSADHMDWPYGYRRLLRFSTQIYNLGRTDFRPKT 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        541 LVQETAYLEDRPLSQLYCAHEENCLSKSADHMDWPYGYRRLLRFSTQIYNLGRTDFRPKT 600

Qy        601 GRDSWVWHQCHRHYHSIEVFTHYDLLTLNGSKVAEGHKASFCLEDTNCPTGLQRRYACAN 660
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        601 GRDSWVWHQCHRHYHSIEVFTHYDLLTLNGSKVAEGHKASFCLEDTNCPTGLQRRYACAN 660

Qy        661 FGEQGVTVGCWDTYRHDIDCQWVDITDVGPGNYIFQVIVNPHYEVAESDFSNNMLQCRCK 720
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db        661 FGEQGVTVGCWDTYRHDIDCQWVDITDVGPGNYIFQVIVNPHYEVAESDFSNNMLQCRCK 720

Qy        721 YDGHRVWLHNCHTGNSYPANAELSLEQEQRLRNNLI 756
              |||||||||||||||||||||||||||||||||||||
Db        721 YDGHRVWLHNCHTGNSYPANAELSLEQEQRLRNNLI 756
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US-09-924-946-2
; Sequence 2, Application US/09924946
; Patent No. US20020102645A1
; GENERAL INFORMATION:
; APPLICANT: American Home Products Corporation
; APPLICANT: Evans, Mark
; APPLICANT: Scicchitano, Marshall
; APPLICANT: Bapat, Ashok
; APPLICANT: Beer, Eric
; APPLICANT: Bhat, Ramesh
; APPLICANT: Ferris, Elissa
; APPLICANT: Mastroeni, Rob
; APPLICANT: Zhang, Jianxiong
; APPLICANT: Karathanasis, Sotirios K.
; TITLE OF INVENTION: A No. US20020102645A1el Member of the Lysyl Oxidase Gene Family
; FILE REFERENCE: 0630/1G703-US2
; CURRENT APPLICATION NUMBER: US/09/924,946
; CURRENT FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: 60/223,763
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/255,838
; PRIOR FILING DATE: 2000-12-15
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 756
; TYPE: PRT
; ORGANISM: Human
US-09-924-946-2

Query Match 100.0%; Score 103; DB 9; Length 756;
Best Local Similarity 100.0%; Pred. No. 4.4e-99;
Matches 103; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 LRLVGPESKPEEGRLEVLHQGQWGTVCDDNFAIQEATVACRQLGFEAALTWAHSAKYGQG 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         32 LRLVGPESKPEEGRLEVLHQGQWGTVCDDNFAIQEATVACRQLGFEAALTWAHSAKYGQG 91

Qy         61 EGPIWLDNVRCVGTESSLDQCGSNGWGVSDCSHSEDVGVICHP 103
              |||||||||||||||||||||||||||||||||||||||||||
Db         92 EGPIWLDNVRCVGTESSLDQCGSNGWGVSDCSHSEDVGVICHP 134
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RESULT 6
US-09-924-946-2
; Sequence 2, Application US/09924946
; Patent No. US20020102645A1
; GENERAL INFORMATION:
; APPLICANT: American Home Products Corporation
; APPLICANT: Evans, Mark
; APPLICANT: Scicchitano, Marshall
; APPLICANT: Bapat, Ashok
; APPLICANT: Beer, Eric
; APPLICANT: Bhat, Ramesh
; APPLICANT: Ferris, Elissa
; APPLICANT: Mastroeni, Rob
; APPLICANT: Zhang, Jianxiong
; APPLICANT: Karathanasis, Sotirios K.
; TITLE OF INVENTION: A No. US20020102645A1el Member of the Lysyl Oxidase Gene Family
; FILE REFERENCE: 0630/1G703-US2
; CURRENT APPLICATION NUMBER: US/09/924,946
; CURRENT FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: 60/223,763
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/255,838
; PRIOR FILING DATE: 2000-12-15
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 756
; TYPE: PRT
; ORGANISM: Human
US-09-924-946-2

Query Match 100.0%; Score 125; DB 9; Length 756;
Best Local Similarity 100.0%; Pred. No. 1.2e-121;
Matches 125; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 PILASAKQHSPVTEGAVEVKYEGHWRQVCDQGWTMNNSRVVCGMLGFPSEVPVDSHYYRK 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        163 PILASAKQHSPVTEGAVEVKYEGHWRQVCDQGWTMNNSRVVCGMLGFPSEVPVDSHYYRK 222

Qy         61 VWDLKMRDPKSRLKSLTNKNSFWIHQVTCLGTEPHMANCQVQVAPARGKLRPACPGGMHA 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        223 VWDLKMRDPKSRLKSLTNKNSFWIHQVTCLGTEPHMANCQVQVAPARGKLRPACPGGMHA 282

Qy        121 VVSCV 125
              |||||
Db        283 VVSCV 287
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5

RESULT 3
US-09-924-946-2
; Sequence 2, Application US/09924946
; Patent No. US20020102645A1
; GENERAL INFORMATION:
; APPLICANT: American Home Products Corporation
; APPLICANT: Evans, Mark
; APPLICANT: Scicchitano, Marshall
; APPLICANT: Bapat, Ashok
; APPLICANT: Beer, Eric
; APPLICANT: Bhat, Ramesh
; APPLICANT: Ferris, Elissa
; APPLICANT: Mastroeni, Rob
; APPLICANT: Zhang, Jianxiong
; APPLICANT: Karathanasis, Sotirios K.
; TITLE OF INVENTION: A No. US20020102645A1el Member of the Lysyl Oxidase Gene Family
; FILE REFERENCE: 0630/1G703-US2
; CURRENT APPLICATION NUMBER: US/09/924,946
; CURRENT FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: 60/223,763
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/255,838
; PRIOR FILING DATE: 2000-12-15
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 756
; TYPE: PRT
; ORGANISM: Human
US-09-924-946-2

Query Match 100.0%; Score 101; DB 9; Length 756;
Best Local Similarity 100.0%; Pred. No. 2.5e-90;
Matches 101; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 6
US-09-924-946-2
; Sequence 2, Application US/09924946
; Patent No. US20020102645A1
; GENERAL INFORMATION:
; APPLICANT: American Home Products Corporation
; APPLICANT: Evans, Mark
; APPLICANT: Scicchitano, Marshall
; APPLICANT: Bapat, Ashok
; APPLICANT: Beer, Eric
; APPLICANT: Bhat, Ramesh
; APPLICANT: Ferris, Elissa
; APPLICANT: Mastroeni, Rob
; APPLICANT: Zhang, Jianxiong
; APPLICANT: Karathanasis, Sotirios K.
; TITLE OF INVENTION: A No. US20020102645A1el Member of the Lysyl Oxidase Gene Family
; FILE REFERENCE: 0630/1G703-US2
; CURRENT APPLICATION NUMBER: US/09/924,946
; CURRENT FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: 60/223,763
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/255,838
; PRIOR FILING DATE: 2000-12-15
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 756
; TYPE: PRT
; ORGANISM: Human
US-09-924-946-2

Query Match 100.0%; Score 109; DB 9; Length 756;
Best Local Similarity 100.0%; Pred. No. 5.8e-99;
Matches 109; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 4
US-09-924-946-2
; Sequence 2, Application US/09924946
; Patent No. US20020102645A1
; GENERAL INFORMATION:
; APPLICANT: American Home Products Corporation
; APPLICANT: Evans, Mark
; APPLICANT: Scicchitano, Marshall
; APPLICANT: Bapat, Ashok
; APPLICANT: Beer, Eric
; APPLICANT: Bhat, Ramesh
; APPLICANT: Ferris, Elissa
; APPLICANT: Mastroeni, Rob
; APPLICANT: Zhang, Jianxiong
; APPLICANT: Karathanasis, Sotirios K.
; TITLE OF INVENTION: A No. US20020102645A1el Member of the Lysyl Oxidase Gene Family
; FILE REFERENCE: 0630/1G703-US2
; CURRENT APPLICATION NUMBER: US/09/924,946
; CURRENT FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: 60/223,763
; PRIOR FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: 60/255,838
; PRIOR FILING DATE: 2000-12-15
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 756
; TYPE: PRT
; ORGANISM: Human
US-09-924-946-2

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Best Local Similarity 100.0%; Pred. No. 2.1e-223;
Matches 227; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 4
AX323479
LOCUS AX323479 2976 bp DNA linear PAT 07-JAN-2002
DEFINITION Sequence 1 from Patent WO0192495.
ACCESSION AX323479
VERSION AX323479.1 GI:18094234
KEYWORDS .
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1
AUTHORS Meyers,R.
TITLE A human lysyl oxidase (47765) and uses thereof
JOURNAL Patent: WO 0192495-A 1 06-DEC-2001;
Millennium Pharmaceuticals, Inc. (US)
FEATURES Location/Qualifiers
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ORIGIN

Query Match 78.7%; Score 2844.4; DB 6; Length 2976;
Best Local Similarity 97.6%; Pred. No. 0;
Matches 2872; Conservative 11; Mismatches 59; Indels 0; Gaps 0;

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Db          95 ATGGCGTGGTCCCCACCAGCCACCCTCTTTCTGTTCCTGCTGCTGCTAGGCCAGCCCCCT 154

Qy         180 CCCAGCAGGCCACAGTCACTGGGCACCACTAAGCTCCGGCTGGTGGGCCCAGAGAGCAAG 239
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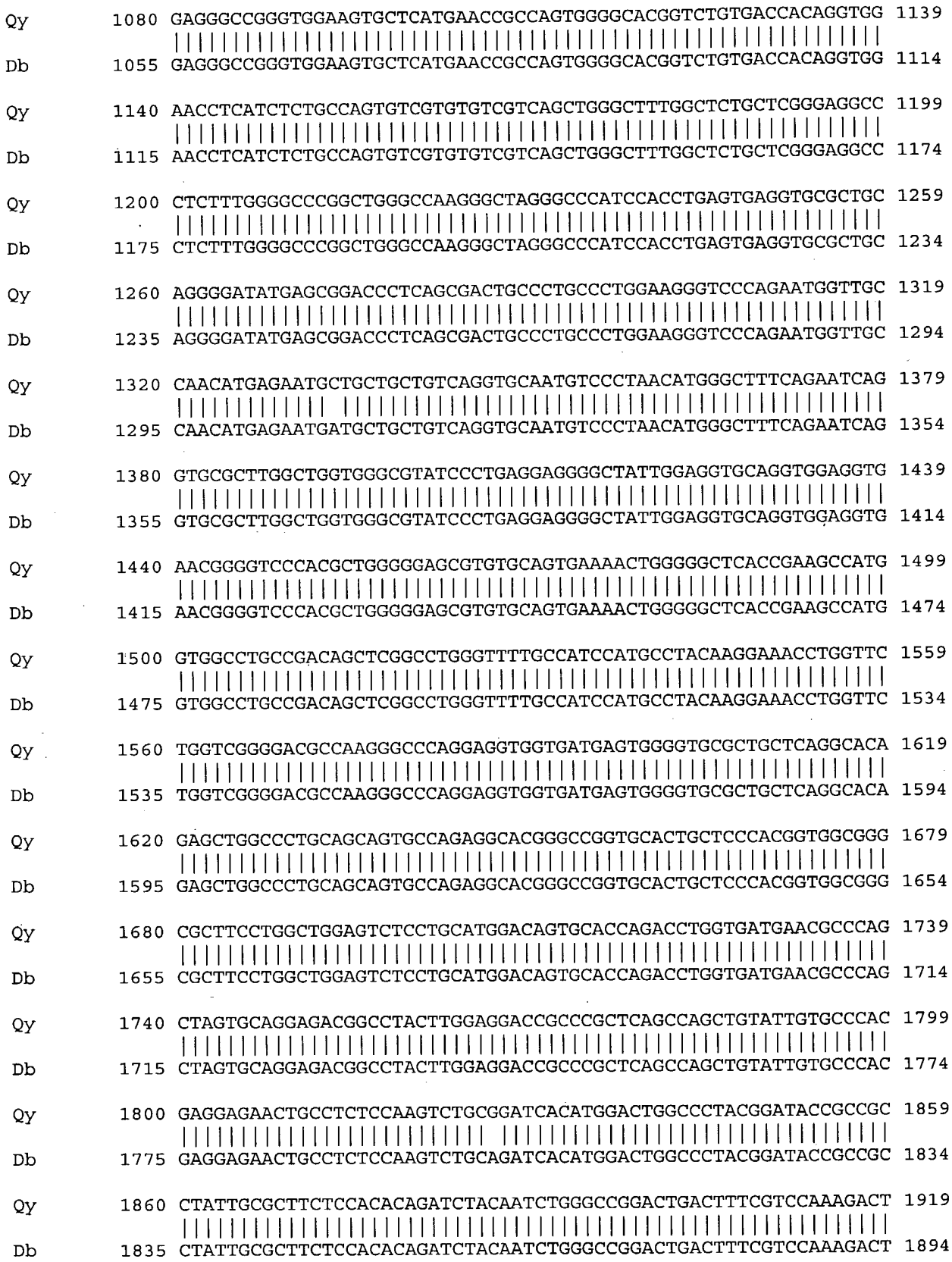
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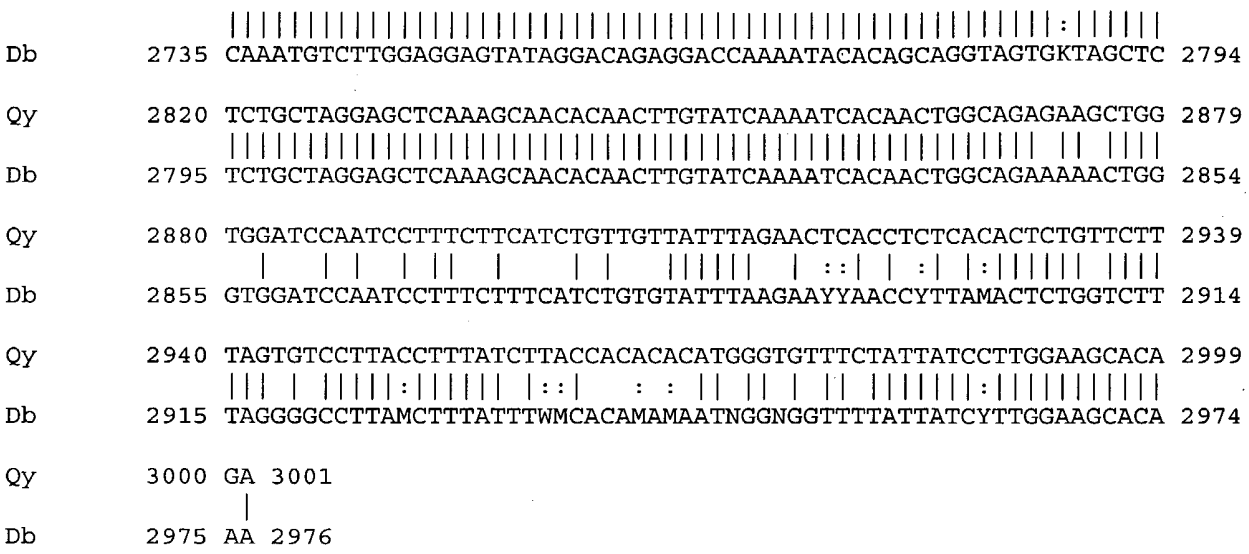
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